

PCT09

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/890,323

DATE: 08/09/2001

TIME: 12:04:10

Input Set : A:\2517-WO Seq.txt

Output Set: N:\CRF3\08092001\I890323.raw

3 <110> APPLICANT: Immunex Corporation
 5 <120> TITLE OF INVENTION: NEW METALLOPROTEINASE-DISINTEGRIN FAMILY MEMBERS: SVPH
 6 DNAS AND POLYPEPTIDES
 8 <130> FILE REFERENCE: 03260.0093-00304
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/890,323
 C--> 11 <141> CURRENT FILING DATE: 2001-07-25
 13 <150> PRIOR APPLICATION NUMBER: 60/116,670
 14 <151> PRIOR FILING DATE: 1999-01-21
 16 <150> PRIOR APPLICATION NUMBER: 60/138,682
 17 <151> PRIOR FILING DATE: 1999-06-14
 19 <150> PRIOR APPLICATION NUMBER: 60/155,798
 20 <151> PRIOR FILING DATE: 1999-09-27
 22 <160> NUMBER OF SEQ ID NOS: 33
 24 <170> SOFTWARE: PatentIn Ver. 2.1
 26 <210> SEQ ID NO: 1
 27 <211> LENGTH: 129
 28 <212> TYPE: DNA
 29 <213> ORGANISM: Homo sapiens
 31 <220> FEATURE:
 32 <223> OTHER INFORMATION: "n" at various positions throughout the sequence
 33 may be any nucleotide
 35 <400> SEQUENCE: 1
 36 atttttgata ccacagtgac caacacgggc acctaagggtg ttcaattcctt tgtagcaagt 60
 W--> 37 ctccacttgca gtattttgcgc ctgcacccaaa aatcctccta cactgttccan ttgcgggtcat 120
 W--> 38 gacangctc 129
 41 <210> SEQ ID NO: 2
 42 <211> LENGTH: 469
 43 <212> TYPE: DNA
 44 <213> ORGANISM: Homo sapiens
 46 <400> SEQUENCE: 2
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 48 caaccacagga ggacatgtga gtcacaatac cctttaatcc acaggttggc tctttggttt 120
 49 ctggaacttt ctgcctcctc taaacqatgt gcgggtgqta cctccctca accagtqat 180
 50 gctttctcac gqgttcaatg aaaaagtctc catgtqtag ttggaataat ccagtcagtc 240
 51 catgqcagtc actgagggct gccgtcccaa ctctggtgcc ctgctqtag accgtgccac 300
 52 taagatggca gaagggggca gaagaaagca tcattttaac atgggagagq ttcccatatc 360
 53 tctttctcat catgtagcta ttggaagaa atccttcatt gacgtccaag ttaaaaaaca 420
 54 gctctctctc ctgctgagaa attctgtagt acaccagtc ctctgagcc 469
 57 <210> SEQ ID NO: 3
 58 <211> LENGTH: 1500
 59 <212> TYPE: DNA
 60 <213> ORGANISM: Homo sapiens
 62 <400> SEQUENCE: 3
 63 caccaggaatt tatattctca aaqaaaatat aatgatgctc ttgcatgctc atttgaataa 60
 64 gctgtttttt tttttttttt tttttttttt tttttttttt tttttttttt tttttttttt

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See page 5

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67 aattgcagtt atatctcttt ttttaaacat atctcttcgg gagcaacatg tctaaataat 300
68 atcccaggac taggttatgt gcttaagaga tgtggaaaca aaattgtgga ggacaatgag 360
69 gaatgtgatt gtqgttcac agaggagtgt cagaaagatc ggtgttgcca atcaaatgtt 420
70 aagttgcaac caggtgccaa ctgtagcatt ggactttgct gtcattgatt tcggtttcgt 480
71 ccattctgat acgtgtgtag gcaggaaagg aatgaatgtg acctgcaga gtaactgcag 540
72 gggaattcaa gttcctgcc aaatgacgtt tataagcagg atggaacccc ttgcaagtat 600
73 gaaggccgtt gtttcaggaa ggggtgcaga tccagatata tgcagtgcc aagcattttt 660
74 ggaacctgat ccattggagg tctagtgaq tctatgatg cagttaactt aataggtgat 720
75 caatttggtt actgtgagat tacagggaatt cgaattttta aaaagtgtga aagtgcacaa 780
76 tcaatatgtg gcaggtctaca gtgtataaat gttgaaacca tccctgattt gccagagcat 840
77 acgaactataa tttctactca tttacaggca gaaaatctca tgtctgtggg cacaggctat 900
78 catctatcca tgaaccccat gggaatacct gacctaggta tgataaatga tggcacctcc 960
79 tgtggagaag gccgggtatg ttttaaaaaa aattgcgtca atagctcagt cctgcagttt 1020
80 gactgtttgc ctgagaaatg caatacccg ggtgttttgc acaacagaaa aaactgccac 1080
81 tgcattgatg ggtggcacc tccattctgt gaggaagtgg ggtatggagg aagcattgac 1140
82 agtgggcttc caggactgct caaqggggcg attccctcgt caatttggtt tgtgtccatc 1200
83 ataattgttc gccattttt attaatcctt tcagtggttt ttgtgtttt ccggcaagtg 1260
84 ataggaaacc acttaaaacc caaacaggaa aaatgccac tatccaaagc aaaaactgaa 1320
85 caggaagaat caaaaacaaa aactgtacag gaagaatcta aaacaaaaac tggacaggaa 1380
86 gaattctgaag caaaaactgg acaggagaa tctaaagcaa aaactggaca ggaagaatct 1440
87 aaagcaaca ttgaaagtaa acgacccaaa gcaaagagtg tcaagaaaca aaaaaagtaa 1500
90 <210> SEQ ID NO: 4
91 <211> LENGTH: 40
92 <212> TYPE: PRT
93 <213> ORGANISM: Homo sapiens
95 <220> FEATURE:
96 <223> OTHER INFORMATION: "Xaa" at various positions throughout the sequence
97 may be any amino acid
99 <400> SEQUENCE: 4

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103 Thr Ala Ser Glu Thr Cys Tyr Lys Glu Leu Asn Thr Leu Gly Asp Arg
104 20 25 30
106 Val Gly His Cys Gly Ile Lys Asn
107 35 40
110 <210> SEQ ID NO: 5
111 <211> LENGTH: 123
112 <212> TYPE: PRT
113 <213> ORGANISM: Homo sapiens
115 <400> SEQUENCE: 5
116 Glu Asp Trp Val Tyr Tyr Arg Ile Ser His Glu Glu Lys Asp Leu Phe
117 1 5 10 15
119 Phe Asn Leu Thr Val Asn Glu Gly Phe Leu Ser Asn Ser Tyr Ile Met
120 20 25 30
122 Glu Lys Arg Tyr Gly Asn Leu Ser His Val Lys Met Met Ala Ser Ser
123 35 40 45
125 Ala Pro Leu Cys His Leu Ser Gly Thr Val Leu Gln Gln Gly Thr Arg
126 50 55 60
128 Val Gly Thr Ala Ala Leu Ser Ala Cys His Glu Thr Thr Thr Thr Thr

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129 65          70          75          80
131 Gln Leu Pro His Gly Asp Phe Phe Ile Glu Pro Val Lys Lys His Pro
132          85          90          95
134 Leu Val Glu Gly Tyr His Pro His Ile Val Tyr Arg Arg Gln Lys
135          100          105          110
137 Val Pro Glu Thr Lys Glu Pro Thr Cys Gly Leu
138          115          120
141 <210> SEQ ID NO: 6
142 <211> LENGTH: 499
143 <212> TYPE: PRT
144 <213> ORGANISM: Homo sapiens
146 <400> SEQUENCE: 6
147 His Glu Asp Leu Tyr Leu Gln Arg Lys Tyr Asn Asp Ala Leu Ala Trp
148 1 5 10 15
150 Ser Phe Gly Lys Val Cys Ser Leu Glu Tyr Ala Gly Ser Val Ser Thr
151 20 25 30
153 Leu Leu Asp Thr Asn Ile Leu Ala Pro Ala Thr Trp Ser Ala His Glu
154 35 40 45
156 Leu Gly His Ala Val Gly Met Ser His Asp Glu Gln Tyr Cys Gln Cys
157 50 55 60
159 Arg Gly Arg Pro Asn Cys Ile Met Gly Ser Gly Arg Thr Gly Phe Ser
160 65 70 75 80
162 Asn Cys Ser Tyr Ile Ser Phe Phe Lys His Ile Ser Ser Gly Ala Thr
163 85 90 95
165 Cys Leu Asn Asn Ile Pro Gly Leu Gly Tyr Val Leu Lys Arg Cys Gly
166 100 105 110
168 Asn Lys Ile Val Glu Asp Asn Glu Glu Cys Asp Cys Gly Ser Thr Glu
169 115 120 125
171 Glu Cys Gln Lys Asp Arg Cys Cys Gln Ser Asn Cys Lys Leu Gln Pro
172 130 135 140
174 Gly Ala Asn Cys Ser Ile Gly Leu Cys Cys His Asp Cys Arg Phe Arg
175 145 150 155 160
177 Pro Ser Gly Tyr Val Cys Arg Gln Glu Gly Asn Glu Cys Asp Leu Ala
178 165 170 175
180 Glu Tyr Cys Asp Gly Asn Ser Ser Ser Cys Pro Asn Asp Val Tyr Lys
181 180 185 190
183 Gln Asp Gly Thr Pro Cys Lys Tyr Glu Gly Arg Cys Phe Arg Lys Gly
184 195 200 205
186 Cys Arg Ser Arg Tyr Met Gln Cys Gln Ser Ile Phe Gly Pro Asp Ala
187 210 215 220
189 Met Glu Ala Pro Ser Glu Cys Tyr Asp Ala Val Asn Leu Ile Gly Asp
190 225 230 235 240
192 Gln Phe Gly Asn Cys Glu Ile Thr Gly Ile Arg Asn Phe Lys Lys Cys
193 245 250 255
195 Glu Ser Ala Asn Ser Ile Cys Gly Arg Leu Gln Cys Ile Asn Val Glu
196 260 265 270
198 Thr Ile Pro Asp Leu Pro Glu His Thr Thr Ile Ile Ser Thr His Leu
199 275 280 285
201 Gln Ala Glu Asn Leu Met Cys Trp Gly Thr Gly Tyr His Leu Ser Pro

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202      290      295      300
204 Lys Pro Met Gly Ile Pro Asp Leu Gly Met Ile Asn Asp Gly Thr Ser
205 305      310      315      320
207 Cys Gly Glu Gly Arg Val Cys Phe Lys Lys Asn Cys Val Asn Ser Ser
208      325      330      335
210 Val Leu Gln Phe Asp Cys Leu Pro Glu Lys Cys Asn Thr Arg Gly Val
211      340      345      350
213 Cys Asn Asn Arg Lys Asn Cys His Cys Met Tyr Gly Trp Ala Pro Pro
214      355      360      365
216 Phe Cys Glu Glu Val Gly Tyr Gly Gly Ser Ile Asp Ser Gly Pro Pro
217      370      375      380
219 Gly Leu Leu Arg Gly Ala Ile Pro Ser Ser Ile Trp Val Val Ser Ile
220 385      390      395      400
222 Ile Met Phe Arg Leu Ile Leu Leu Ile Leu Ser Val Val Phe Val Phe
223      405      410      415
225 Phe Arg Gln Val Ile Gly Asn His Leu Lys Pro Lys Gln Glu Lys Met
226      420      425      430
228 Pro Leu Ser Lys Ala Lys Thr Glu Gln Glu Glu Ser Lys Thr Lys Thr
229      435      440      445
231 Val Gln Glu Glu Ser Lys Thr Lys Thr Gly Gln Glu Glu Ser Glu Ala
232      450      455      460
234 Lys Thr Gly Gln Glu Glu Ser Lys Ala Lys Thr Gly Gln Glu Glu Ser
235 465      470      475      480
237 Lys Ala Asn Ile Glu Ser Lys Arg Pro Lys Ala Lys Ser Val Lys Lys
238      485      490      495
240 Gln Lys Lys
244 <210> SEQ ID NO: 7
245 <211> LENGTH: 2301
246 <212> TYPE: DNA
247 <213> ORGANISM: Homo sapiens
249 <400> SEQUENCE: 7
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251 caggatgagc acccccaata tcacagccct ccggatgtgg tgattcctgt gaggataact 120
252 ggcaccacca gaggcattgac acctccaggc tggctctcct atatcctgcc ctttggaggc 180
253 cagaaacaca ttatccacat aaaggtaag agcttttgt ttccaaaca cctccctgtg 240
254 ttacactaca cagaccaggg tgcatacctt gaggaccagc catttgtcca gaataactgc 300
255 tactatcatg gttatgtgga aggggaccca gaatccctgg ttccctcag taactgtttt 360
256 qggggttttc aaggaatatt acagataaat gactttgctt atqaaatcaa qccctcagca 420
257 ttttctacca cgtttgaaca tetggtatcc aagatggaca gtgaggagaa acatttttca 480
258 accatgagat ccggaattat gcaaaatgaa ataactgccc qaatggaatt tgaagaaatt 540
259 gataattcca ctcaagaaga aagttcttat gtggctggtt qgattccatt taggattggt 600
260 gaaattgtag tcttcattga taattatctg tacattcgtt atgaaaggaa cgaatcaaaq 660
261 ttgctggagg atctatatgt tattgttaat atagtggatt ccattttgga tctcattggt 720
262 gtttaagggtg tattattttg tttggagatc tggaccaata aaaacctcat tctagttagt 780
263 gatgtaagga aatctgtgca cctgtattgc aagtggaggt cggagaacat taqccccgg 840
264 atgcaacatg acacctcaca tctttcaca actctaggat taagagggtt aagtggcata 900
265 gtagctttta gaqqaatgtg tacaccacac ctaqgttgtg caattgttac tttcatqaac 960
266 aaaactttgq gcacttttcc aattgcagtg gctcatcacc taqgtcataa tttgggcatg 1020
267 aacctgatag aagatacata tctttattca caacctagat gcaattatc tttttttttt 1080

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268 ccaccaataa ctaaatttag caattgtagt tatggtgatt tttgggaata tactgtagag 1140
269 aggacaaagt gtttgcttga aacagtacac acaaaggaca tctttaatgt gaagcgtgt 1200
270 gggaaatggtg ttgttgaaga aggagaagag tgtgactgtg gaccttfaaa gcattgtgca 1260
271 aaagatccct gctgtctgtc aaattgcact ctgactgatg gttctacttg tgcctttggg 1320
272 ctttgttgca aagactgcaa gttcctacca tcagggaag tgtgtagaaa gtaggtcaat 1380
273 gaatgtgate ttccagagtg gtgcaatggt acttccata agtgcccaga tgaactttat 1440
274 gtggaagatg gaattccctg taaggagagg gctactgct atgaaaagag ctgtcatgac 1500
275 cgcaatgaac agtgtaggag gatttttggg gcaggcgcaa atactgcaag tgagacttgc 1560
276 taaaaagaat tgaacacctt aggtgacctt gttggctact gtggtatcaa aaatgctaca 1620
277 tatataaagt gtaatatctc agatgtccag tgtggaagaa ttcagtgtga gaatgtgaca 1680
278 gaaattccca atatgagtga tcatactact gtgcattggg ctgccttcaa tgacataatg 1740
279 tgcctggagta ctgattacca tttggggatg aagggacctg atattggtga agtgaaagat 1800
280 ggaacagagt gtgggataga tcatatatgc atccacaggc actgtgtcca tataaccatc 1860
281 ttgaatagta attgctcacc tgcattttgt aacaagaggg gcactctgca caataaacat 1920
282 cactgccatt gcaattatct gtgggacctt cccaactgcc tgataaaagg ctatggaggt 1980
283 agtggttaca gtggcccacc cctaagaga aagaagaaaa aqaagttctg tctctgtgt 2040
284 atattgttgc liattgtttt gtttatttta ttatgttgtc ttatcgact ttgtaaaaaa 2100
285 agtaaaccaa taaaaagca gcaagatgtt caaactccat ctgcaaaaga agaggaaaaa 2160
286 attcagctc gacctcatga gttacctccc cagagtcaac cttgggtgat gcttcccag 2220
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292 <211> LENGTH: 2364
293 <212> TYPE: DNA
294 <213> ORGANISM: Homo sapiens
296 <400> SEQUENCE: 8
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298 caggatgagc acccccataa tcacagccct ccgatgtgg tgattcctgt gaggataact 120
299 ggcaccacca gaggcattga acctccaggc tggctctcct atactctgcc ctttggaggc 180
300 cagaaacaca ttatccacat aaaggtaag agcttttgt ttccaaaca cctccctgtg 240
301 ttacactaca cagaccaggg tgcatacctt gaggaccagc catttgctca gaataactgc 300
302 tactatcatg gttatgtgga aggggaccca gaatccctgg ttccctcag tacctgtttt 360
303 gggggttttc aaggaatatt acagataaat gactttgctt atgaaatcaa gcccttagca 420
304 ttttctacca cgtttgaaca tctggtatac aagatggaca gtgaggagaa acaattttca 480
305 accatgagat ccggttttat gcaaaatgaa ataactgcc gaatggaatt tgaagaaatt 540
306 gataattcca ctccagaaga aagttcttat gtgggctggt ggaatccatt taggattgtt 600
307 gaaattgtag tcgtcattga taattatctg tacattcgtt atgaaaggaa cgaactcaag 660
308 ttgctggagg atctatatgt tattgttaat atagtggatt ccattttgga tgtcatttgt 720
309 gtttaaggtg tattattttg tttggagatc ttgaaccaaa aaaacctcat tgtagttagt 780
310 gatgtaagga aatctgtgca cctgtattgc aagtggaggt cggagaacat tacgcccagg 840
311 atgcaacatg acaactcaca tcttttcaca actctaggat taagaaggtt aagtggcata 900
312 ggagctttta gaggaaatgt tacaccacac cgtagttgtg caattgttac ttatcatgaa 960
313 aaaacttttg gcacttttct aattgcagtg gctcctcctc taggtcataa ttggggcatg 1020
314 aaccatgatg aggatacatg tcgttgttca caacctagat gcataatgca tgaaggcaac 1080
315 ccaccaataa ctaaatttag caattgtagt tatggtgatt tttgggaata tactgtagag 1140
316 aggacaaagt gtttgcttga aacagtacac acaaaggaca tctttaatgt gaagcgtgt 1200
317 gggaaatggtg ttgttgaaga aggagaagag tgtgactgtg gaccttfaaa gcattgtgca 1260
318 aaagatccct gctgtctgtc aaattgcact ctgactgatg gttctacttg tgcctttggg 1320
319 ctttgttgca aagactgcaa gttcctacca tcagggaag tgtgtagaaa gtaggtcaat 1380

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Sequence data were generated using the following protocol:
 ensure a minimum of 100 bp of sequence is present
 in the <220> to <223> fields of each sequence
 using n or Xaa

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/890,323

DATE: 08/09/2001

TIME: 12:04:11

Input Set : A:\2517-WO Seq.txt

Output Set: N:\CRF3\08092001\I890323.raw

L:10 M:270 C: Current Application Number differs, Replaced Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:37 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:1
L:37 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:1
L:37 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:38 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:1
L:38 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:1
L:38 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
L:100 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:4
L:100 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:4
L:100 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:1471 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:31
L:1471 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:31
L:1471 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:1502 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:33
L:1502 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:33
L:1502 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33